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				Application Number	Filed concurrently herewith
				Filing Date	September 28, 2001
				First Named Inventor	Eric C. Hannah
				Group Art Unit	Not yet assigned
				Examiner Name	Not yet assigned
				Attorney Docket Number	42390P11816
Sheet	1	of	2		

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FOREIGN PATENT DOCUMENTS								
Examiner Initials*	Cite No. <sup>1</sup>	Office <sup>3</sup>	Foreign Patent Number <sup>4</sup>	Kind Code <sup>5</sup> (if known)	Name of Patentee or Applicant of Cited Document	Date of Publication of Cited Document MM-DD-YYYY	Pages, Columns, Lines Where Relevant Passages or Relevant Figures Appear	T <sup>6</sup>
			WO 98/48270		California Institute of Technology	10-29-1998		

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**OTHER PRIOR ART - NON PATENT LITERATURE DOCUMENTS**

Examiner Initials	Cite No. <sup>1</sup>	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date page(s), volume-issue number(s), publisher, city and/or country where published.	T <sup>2</sup>
		Al-Lazikani, Bissan <i>et al.</i> , "Protein structure prediction", <i>Curr. Opin. in Chem. Biol.</i> , 5:51-56, 2001	
		Bonneau and Baker, "Ab Initio Protein Structure Prediction: Progress and Prospects", <i>Annu. Rev. Biophys. Biomol. Struct.</i> , 30:73-189, 2001	
		Clementi, Cecilia <i>et al.</i> , "Topological and Energetic Factors: What Determines the Structural Details of the Transition State Ensemble and "En-route" Intermediates for Protein Folding? An Investigation for Small Globular Proteins", <i>J. Mol. Biol.</i> , 298:937-953, 2000	
		Debe, Derek A. <i>et al.</i> , "The topomer-sampling model of protein folding", <i>Proc. Natl. Acad. Sci USA</i> , 96:2596-2601, March 1999	
		Debe and Goddard III, "First Principles Prediction of Protein Folding Rates", <i>J. Mol. Biol.</i> , 294:619-625, 1999	
		Eyrich, Volker A. <i>et al.</i> , "Prediction of Protein Tertiary Structure to Low Resolution: Performance for a Large and Structurally Diverse Test Set", <i>J. Mol. Biol.</i> 288:725-742, 1999	
		Mirny, Leonid A. <i>et al.</i> , "Statistical significance of protein structure prediction by threading", <i>PNAS</i> , 97(18) 9978-9983, August 29, 2000	
		Pillardy, Jaroslaw <i>et al.</i> , "Recent improvements in prediction of protein structure by global optimization of a potential energy function", <i>PNAS</i> 98(5) 2329-2333, February 27, 2001	
		Thomas and Dill, "An iterative method for extracting energy-like quantities from protein structures", <i>Proc. Natl. Acad. Sci USA</i> 93:11628-11633, October 1996	

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